

## Transformation of 5-Hydroxy- to (5-Chloropentanoyl)amino Derivatives under ‘Direct Amide Cyclization’ Conditions

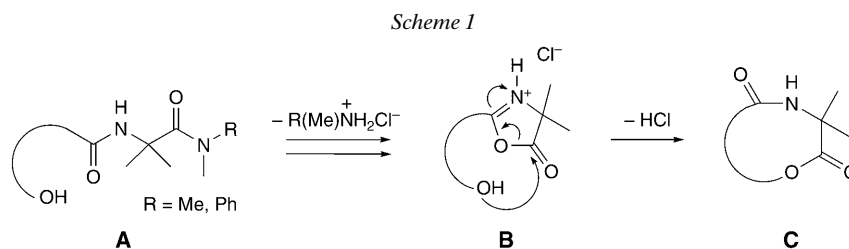
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The application of the ‘direct amide cyclization’ conditions to the linear  $\delta$ -hydroxy diamide **11** is described (Scheme 3). Instead of the cyclization to the expected nine-membered cyclodepsipeptide, only the chloro acid **12** was obtained. Its formation could be explained by consecutive formation of the 1,3-oxazol-5(4*H*)-one **16** and the six-membered imino lactone **17** as intermediates (Scheme 4). The spontaneous isomerization of the latter gave **12** in a good yield.

**1. Introduction.** – Cyclic depsipeptides are a class of biologically active secondary metabolites, which contain not only amide bonds as part of their ring structure, but also other, usually lactone, bonds. Their biological activity is diverse, being based mainly on their capability of selectively transporting metal ions through cell membranes, and they are sought after as potential anticancer, antiviral, antibiotic, and anti-inflammatory drugs [1].

One of the many methods known for their synthesis is the direct amide cyclization [2]: a suspension of an amide of type **A** in toluene is treated with dry HCl gas. Cyclization by elimination of the corresponding ammonium chloride leads to the intermediate 1,3-oxazol-5(4*H*)-one of type **B**. In the absence of other nucleophiles, **B** undergoes a ring enlargement *via* intramolecular nucleophilic attack of the OH group at the carbonyl C-atom of the neighboring lactone group, to give the cyclodepsipeptide **C** (Scheme 1).

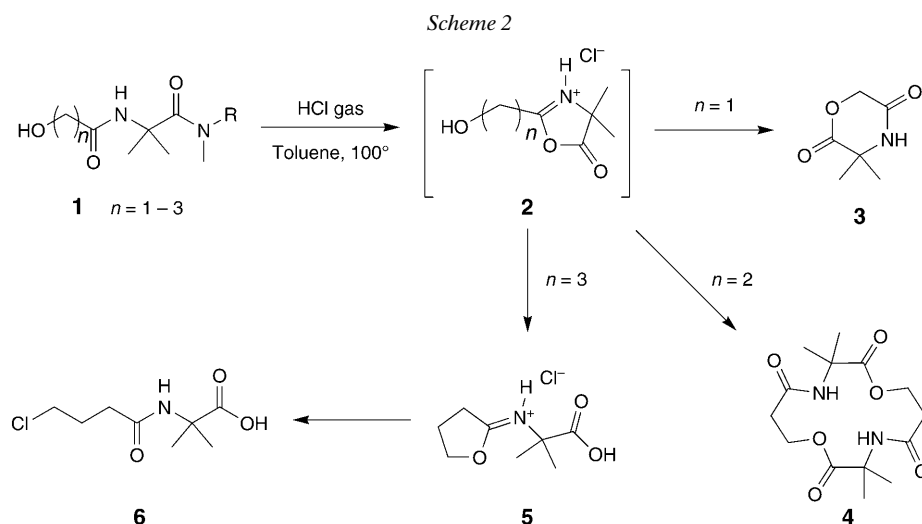


This method has been used successfully for the synthesis of 6-, 9-, 12-, and 15-membered [3], and larger ring systems [4]. For example,  $\alpha$ -hydroxy acid derivatives of type **1**

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( $n=1$ ) under the ‘direct amide cyclization’ conditions led to morpholinediones of type **3** [5] (Scheme 2). Recently, we have shown that diamides of type **A**, which contain  $\beta$ -hydroxy acid moieties, *i.e.*, **1** ( $n=2$ ), under these reaction conditions, did not give the expected seven-membered cyclodepsipeptides, and only their dimers, the 14-membered rings **4**, were isolated in good yields [6] [7] (Scheme 2). The treatment of  $\gamma$ -hydroxy acid derivatives (**1**,  $n=3$ ) with HCl gas in toluene yielded no cyclodepsipeptides at all. Instead, the only products isolated were the hydrochlorides of imino lactones **5**, which are unstable in solution and isomerize in polar solvents or on silica gel to give the chlorinated acids **6** [8] (Scheme 2).



In all cases so far, the formation of the intermediate oxazolone **2** was established chemically or by IR spectroscopy. The results show that there is a relationship between the number of C-atoms between the OH and CONH groups and the type of product formed. Therefore, it was of interest to subject  $\delta$ -hydroxy amides **1** ( $n=4$ ) to the conditions of the ‘direct amide cyclization’, which could yield either a 9- or 18-membered cyclodepsipeptide on the one hand, or a six-membered imino lactone on the other. The result of a first example is shown below.

**2. Results and Discussion.** – Since the previously used standard methods for syntheses of  $\beta$ - and  $\gamma$ -hydroxy acids [8] were not applicable to the  $\delta$ -hydroxy analogues, we used the method of *Goto et al.* [9]: 3,3-dimethyl- $\gamma$ -pentano-5-lactone (**7**) was treated with NaOH and  $\text{Ac}_2\text{O}$ , which led to the *O*-protected 5-acetoxy-3,3-dimethylpentanoic acid (**8**) in moderate yield (Scheme 3).

The reaction of the acid **8** with 2,2,*N*-trimethyl-*N*-phenyl-2*H*-azirin-3-amine (**9**) [10] yielded the protected diamide **10**. After deprotection with LiOH, **11** was subjected to the ‘direct amide cyclization’ (DAC) conditions (HCl gas, toluene,  $100^\circ$ ). Trituration of the crude product with  $\text{CH}_2\text{Cl}_2$ , a procedure that has allowed the isolation of the imino lactone hydrochlorides **5** without isomerization to the corresponding chlorinated acids [8], yielded in the present case the chloro acid **12** directly as the only product in good

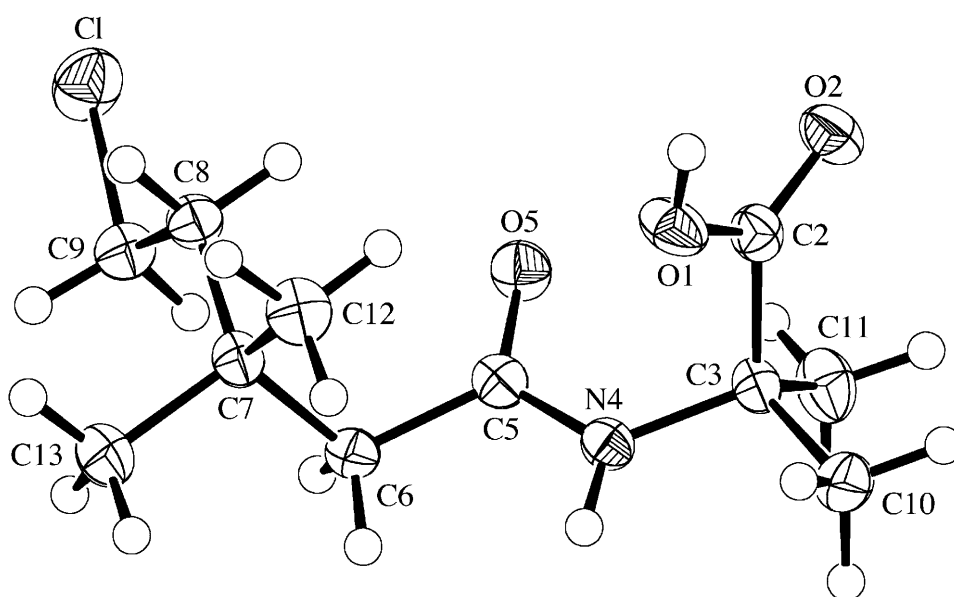
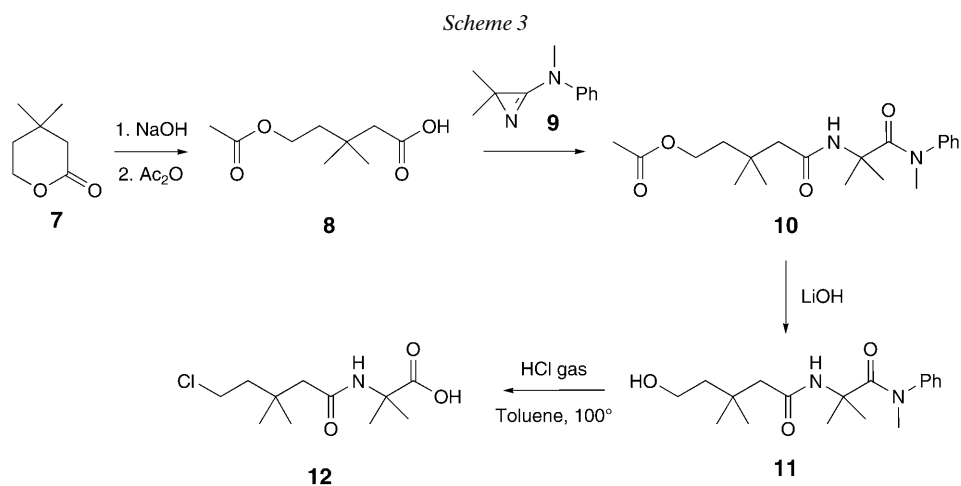


Fig. 1. ORTEP Plot [11] of the molecular structure of **12**. Arbitrary numbering of the atoms; 50% probability ellipsoids.

yield. Recrystallization from MeCN gave crystals which were suitable for an X-ray crystal-structure determination (Fig. 1).

Although the compound is achiral, it has crystallized in a polar space group and the absolute structure has been determined unambiguously. The OH group forms an intermolecular H-bond with the amide O-atom of a neighboring molecule, thereby linking the molecules into extended chains, which run parallel to the [0 1 0] direction and can be described by a graph set motif [12] of C(7). The amide group forms an intermolec-

ular H-bond with the carboxy carbonyl O-atom of a different neighboring molecule. This interaction links the molecules into extended chains, which run parallel to the  $[1\ 0\ 0]$  direction and can be described by a graph set motif of C(5). The combination of both interactions generates a two-dimensional framework that lies parallel to the  $(0\ 0\ 1)$  plane (Fig. 2).

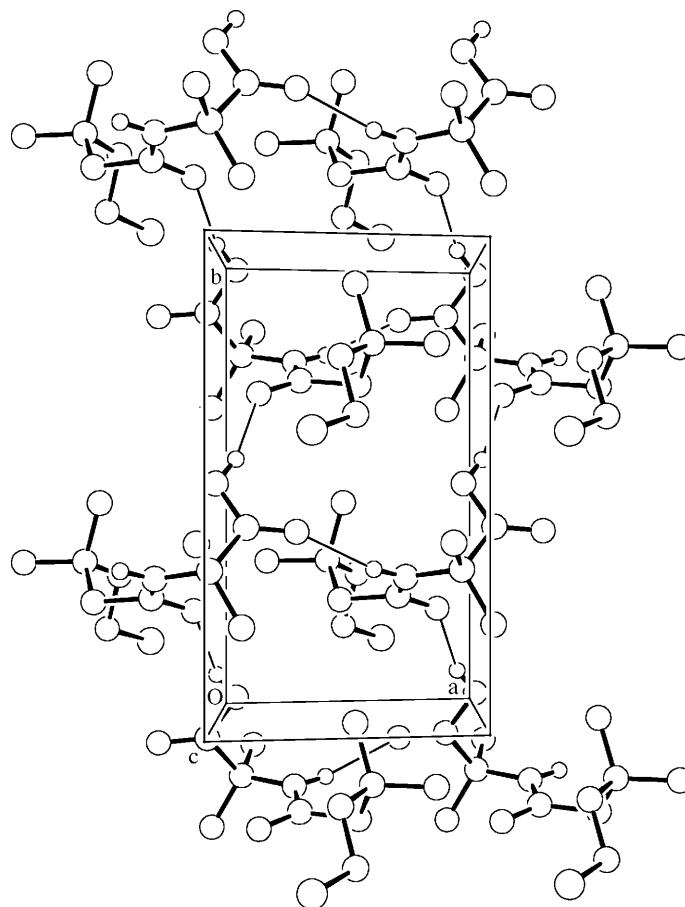
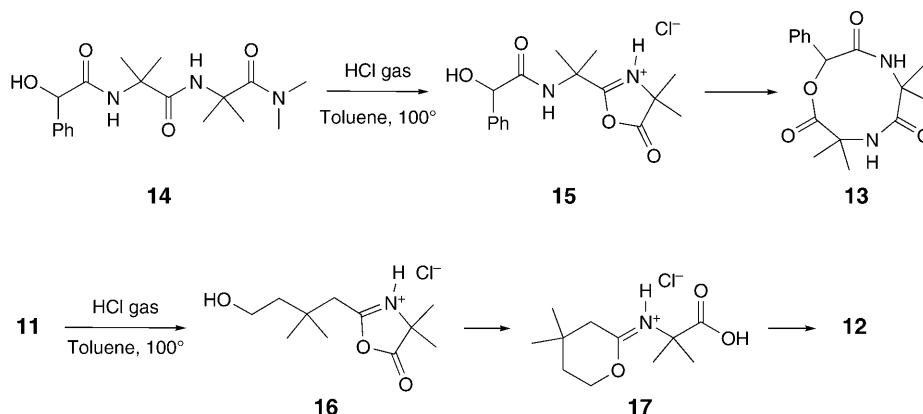


Fig. 2. Molecular packing of **12**, showing the H-bonding interactions

In our earlier studies, it has been observed that the ring size has a controlling influence on the formation of possible products from compounds of type **1** under the DAC conditions. With  $\beta$ -hydroxy acids ( $n=2$ , Scheme 2), fourteen-membered rings **4** (cyclodimers) are preferred over seven-membered ones, while the formation of five-membered imino lactones **5** is preferred over the corresponding eight-membered cyclodepsipeptides in the case of  $\gamma$ -hydroxy acids ( $n=3$ ). Therefore, the expected product in the case where **1** contains a  $\delta$ -hydroxy acid moiety was either a six-membered imino lactone analogous to **5**, or a nine-membered cyclodepsipeptide. As the nine-membered

Scheme 4



cyclodepsipeptide **13** has already been synthesized *via* DAC from the linear precursor **14** [3][13] (Scheme 4), the formation of **12** was a surprise.

The formation of **12** proceeds most probably through the intermediate 1,3-oxazolone derivative **16** and the six-membered imino lactone **17**, which apparently is unstable under these conditions and, by analogy with **5**, isomerizes spontaneously to give the chloro acid **12**.

**3. Conclusions.** – The DAC conditions were applied to the linear precursor **11**, which contains a  $\delta$ -hydroxy acid moiety. Surprisingly, no cyclic depsipeptide was formed under these conditions. Thus, the dependence of the result of the DAC reaction on whether an  $\alpha$ -,  $\beta$ -,  $\gamma$ -, or  $\delta$ -hydroxy-acid-moiety-containing diamide is used has been confirmed. The general use of the DAC for the synthesis of cyclic depsipeptides is apparently limited to linear precursors which contain  $\alpha$ -hydroxy-acid-moiety-containing amides. The use of  $\beta$ -hydroxy acid precursors is also possible for the formation of larger rings [4], but can lead to cyclodimers or side products as well.

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### Experimental Part

1. *General.* See [8].

2. *Starting Materials.* 2,2-Trimethyl-N-phenyl-2H-azirin-3-amine (**9**) was prepared according to standard procedures (*cf.* [8] and refs. cit. therein). Pentanolactone **7** was prepared by a known method [14].

3. *5-Acetoxy-3,3-dimethylpentanoic Acid (8).* To a soln. of **7** (20 mmol, 2560 mg) in MeOH (10 ml) was added 2N NaOH (11 ml) at 0°. The mixture was stirred at r.t. for 2 h, the solvent was evaporated *i.v.*, and the remaining H<sub>2</sub>O was distilled azeotropically with benzene (3 × 10 ml). The white residue was dried overnight under h.v., and then Ac<sub>2</sub>O (10 ml) was added. After 14 h at 80°, the mixture was cooled, the solvent evaporated, and the oily residue was extracted with AcOEt (5 × 30 ml). Drying (MgSO<sub>4</sub>) and CC (SiO<sub>2</sub>; CH<sub>2</sub>Cl<sub>2</sub>/MeOH 20 : 1) yielded 1201 mg (32%) of **9**. Recovered starting material:

980 mg (37%). IR: 3288s (br.), 2971s, 1738vs, 1711vs, 1471w, 1378s, 1356m, 1244s, 1094s, 1040s, 925w. <sup>1</sup>H-NMR: 0.97 (s, Me<sub>2</sub>C); 1.59–1.71 (m, CH<sub>2</sub>); 1.99 (s, MeCO); 2.21 (s, CH<sub>2</sub>); 4.01 (br. s, CH<sub>2</sub>O); 10.41 (br. s, COOH). <sup>13</sup>C-NMR: 20.7 (q, MeCO); 27.2 (q, Me<sub>2</sub>C); 32.0 (s, Me<sub>2</sub>C); 37.6, 43.9 (2t, 2 CH<sub>2</sub>); 66.6 (t, CH<sub>2</sub>O); 171.1 (s, C=O); 177.3 (s, COOH). ESI-MS: 211 (100, [M+Na]<sup>+</sup>).

4. 4-[1-Methyl-1-(N-methyl-N-phenylcarbamoyl)ethylcarbamoyl]-3,3-dimethylbutyl Acetate (**10**). To a soln. of **8** (376 mg, 2 mmol) in dry THF (20 ml), **9** (365 mg, 2.1 mmol) was added. The mixture was stirred at r.t. overnight, and the solvent was evaporated *i.v.*, CC (CH<sub>2</sub>Cl<sub>2</sub>/MeOH 40:1) yielded 622 mg (86%) of **10**. Colorless powder. M.p. 98.1–99.4°. <sup>1</sup>H-NMR: 0.97, 1.41 (2s, 2 Me<sub>2</sub>C); 1.62 (t, *J* = 7.1, CH<sub>2</sub>); 1.99 (s, MeCO); 2.08 (s, CH<sub>2</sub>); 3.19 (s, MeN); 3.10 (t, *J* = 7.1, CH<sub>2</sub>O); 6.55 (s, NH); 7.20–7.39 (m, 5 arom. H). <sup>13</sup>C-NMR: 20.8 (q, MeCO); 26.3, 27.0 (2q, 2 Me<sub>2</sub>C); 33.0 (s, Me<sub>2</sub>C); 41.3 (t, CH<sub>2</sub>); 42.3 (q, MeN); 48.2 (t, CH<sub>2</sub>); 58.1 (s, Me<sub>2</sub>C); 68.7 (t, CH<sub>2</sub>O); 127.8, 128.0, 129.3 (3d, 5 arom. CH); 144.6 (s, arom. C); 169.8, 171.6, 173.2 (3s, 3 C=O). ESI-MS: 385 (100, [M+Na]<sup>+</sup>).

5. 5-Hydroxy-3,3-dimethyl-N-[1-methyl-1-(N-methyl-N-phenylcarbamoyl)ethyl]pentanamide (**11**). A soln. of **9** (362 mg, 1 mmol) in THF/H<sub>2</sub>O 2:1 (20 ml) was treated with 4 equiv. of LiOH at r.t. for 4 h. Evaporation of the solvent *i.v.*, extraction of the residue with CH<sub>2</sub>Cl<sub>2</sub>, drying (MgSO<sub>4</sub>), evaporation *i.v.*, and washing with Et<sub>2</sub>O yielded **11**, which was used without further purification. Yield: 298 mg (93%) of **11**. White solid. M.p. 128.4–126.0°. <sup>1</sup>H-NMR: 0.99, 1.46 (2s, 2 Me<sub>2</sub>C); 1.60 (t, *J* = 7.1, CH<sub>2</sub>); 2.00 (s, CH<sub>2</sub>); 3.17 (s, MeN); 3.18 (t, *J* = 7.1, CH<sub>2</sub>O); 6.51 (s, NH); 7.18–7.36 (m, 5 arom. H). <sup>13</sup>C-NMR: 26.4, 28.8 (2q, 2 Me<sub>2</sub>C); 32.8 (s, Me<sub>2</sub>C); 41.4 (t, CH<sub>2</sub>); 42.4 (q, MeN); 48.2 (t, CH<sub>2</sub>); 58.2 (s, Me<sub>2</sub>C); 69.7 (t, CH<sub>2</sub>O); 127.8, 128.0, 129.3 (3d, 5 arom. CH); 144.5 (s, arom. C); 171.7, 173.3 (2s, 2 C=O). ESI-MS: 343 (100, [M+Na]<sup>+</sup>).

6. 2-[5-Chloro-3,3-dimethylpentanoyl]amino]-2-methylpropanoic Acid (**12**). A suspension of **11** (80 mg, 0.25 mmol) in dry toluene (30 ml) was heated to 100°, and HCl gas was bubbled through the suspension for 4–6 min. Then, the mixture was allowed to cool to r.t. while bubbling N<sub>2</sub> through it (*ca.* 20 min). The solvent was evaporated, and the white residue was washed with CH<sub>2</sub>Cl<sub>2</sub> (3 × 15 ml) and dried *in vacuo*: 36 mg (59%) of **12**. M.p. 118.9–121.0°. IR: 3320vs, 2980s, 1722vs, 1620s, 1561s, 1466s, 1428s, 1389m, 1246m, 1232m, 1166s, 1092m, 1051w, 945w. <sup>1</sup>H-NMR ((D<sub>6</sub>)DMSO): 0.97, 1.33 (2s, 2 Me<sub>2</sub>C); 1.68–1.79 (m, CH<sub>2</sub>); 1.97 (s, CH<sub>2</sub>); 3.60–3.69 (m, CH<sub>2</sub>Cl); 7.82 (br. s, NH); 11.81 (br. s, COOH). <sup>13</sup>C-NMR ((D<sub>6</sub>)DMSO): 24.8, 27.2 (2q, 2 Me<sub>2</sub>C); 33.4, 42.5 (2t, 2 CH<sub>2</sub>); 54.6 (t, CH<sub>2</sub>Cl); 55.7 (s, Me<sub>2</sub>C); 169.9, 175.4 (2s, 2 C=O). ESI-MS: 274 (25, [M(<sup>37</sup>Cl)+Na]<sup>+</sup>), 272 (100, [M(<sup>35</sup>Cl)+Na]<sup>+</sup>), 214 (10, [M-Cl]<sup>+</sup>).

7. X-Ray Crystal-Structure Determination of **12** (Table and Figs. 1 and 2)<sup>3</sup>. All measurements were performed on a Nonius KappaCCD area-detector diffractometer [15] using graphite-monochromated MoK<sub>α</sub> radiation (λ 0.71073 Å) and an Oxford Cryosystems Cryostream 700 cooler. The data collection and refinement parameters are given in the Table, and views of the molecules are shown in Figs. 1 and 2. Data reduction was performed with HKL Denzo and Scalepack [16]. The intensities were corrected for Lorentz and polarization effects, and an absorption correction based on the multi-scan method [17] was applied. Equivalent reflections, other than Friedel pairs, were merged. The structure was solved by direct methods using SIR92 [18], which revealed the positions of all non-H-atoms. The non-H-atoms were refined anisotropically. The OH and amide H-atoms were placed in the positions indicated by a difference electron-density map, and their positions were allowed to refine together with individual isotropic displacement parameters. All remaining H-atoms were placed in geometrically calculated positions and refined using a riding model where each H-atom was assigned a fixed isotropic displacement parameter with a value equal to 1.2U<sub>eq</sub> of its parent C-atom (1.5U<sub>eq</sub> for the Me groups). Refinement of the structure was carried out on F<sup>2</sup> using full-matrix least-squares procedures, which minimized the function Σw(F<sub>o</sub><sup>2</sup> - F<sub>c</sub><sup>2</sup>)<sup>2</sup>. A correction for secondary extinction was applied. Refinement of the absolute structure parameter [19][20] yielded a value of 0.01(6), which confirms that the refined model represents the true absolute structure. Neutral atom scattering factors for non-H-atoms were taken from [21], and

<sup>3</sup>) CCDC-286116 contains supplementary crystallographic data for this paper. These data can be obtained free of charge from the Cambridge Crystallographic Data Centre (CCDC), via [http://www.ccdc.cam.ac.uk/data\\_request/cif](http://www.ccdc.cam.ac.uk/data_request/cif).

Table. Crystallographic Data of **12**

Crystallized from	MeCN	$D_x$ [g cm <sup>-3</sup> ]	1.273
Empirical formula	C <sub>11</sub> H <sub>20</sub> ClNO <sub>3</sub>	$\mu$ (MoK $\alpha$ ) [mm <sup>-1</sup> ]	0.286
Formula weight [g mol <sup>-1</sup> ]	249.74	Scan type	$\phi$ and $\omega$
Crystal color, habit	colorless, plate	$2\theta_{(\max)}$ [°]	55
Crystal dimensions [mm]	0.02 × 0.20 × 0.25	Transmission factors (min; max)	0.864; 0.996
Temp. [K]	160(1)	Total reflections measured	13304
Crystal system	monoclinic	Symmetry independent reflections	2950
Space group	$P2_1$	Reflections with $I > 2\sigma(I)$	2548
$Z$	2	Reflections used in refinement	2950
Reflections for cell determination	45269	Parameters refined; restraints	159; 1
$2\theta$ Range for cell determination [°]	4–55	$R$ (on $F$ ; $I > 2\sigma(I)$ reflections)	0.0372
Unit cell parameters	$a$ [Å] 6.1314(3)	$wR$ (on $F^2$ ; all indept. reflections)	0.0814
	$b$ [Å] 10.4871(5)	Weighting parameters [ $a$ ; $b$ ] <sup>a</sup> )	0.0314; 0.1588
	$c$ [Å] 10.4750(5)	Goodness-of-fit	1.044
	$\beta$ [°] 104.760(2)	Secondary extinction coefficient	0.035(5)
	$V$ [Å <sup>3</sup> ] 651.32(5)	Final $\Delta_{\max}/\sigma$	0.001
		$\Delta\rho$ (max; min) [e Å <sup>-3</sup> ]	0.19; –0.16

<sup>a</sup>)  $w^{-1} = \sigma^2(F_o^2) + (aP)^2 + bP$  where  $P = (F_o^2 + 2F_c^2)/3$

the scattering factors for H-atoms were taken from [22]. Anomalous dispersion effects were included in  $F_c$  [23]; the values for  $f'$  and  $f''$  were those of [24]. The values of the mass attenuation coefficients are those of [25]. All calculations were performed using the SHELXL97 program [26].

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